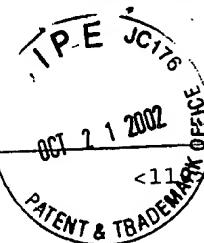


RECEIVED

OCT 24 2002

TECH CENTER 1600/2900



<110> Vogels, Ronald
Schouten, Govert J.
Bout, Abraham

<120> Means and Methods for Fibroblast-Like or Macrophage-Like Cell Transduction

<130> 2183-3982.2US

<140> 09/517,898

<141> 2000-03-03

<150> 60/122,732

<151> 1999-03-03

<160> 38

<170> PatentIn version 3.1

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ber protein derived from adenovirus serotype

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<212> DNA
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<220>
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ber protein derived from adenovirus serotype

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ber protein derived from adenovirus serotype

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<212> DNA
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Cont
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ber protein derived from adenovirus serotype

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37

<210> 31
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ber protein derived from adenovirus serotype

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30

<210> 32
<211> 1068

Ed
int
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<213> Artificial sequence

<220>
<223> DNA encoding Adenovirus Ad5/fib16 chimeric fiber

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agcccagatg gagttctaac tcttaatgt gttaatccac tcactaccgc cagcggaccc	240
ctccaactta aagttggaag cagtcttaca gtagatacta tcgatgggtc tttggaggaa	300
aatataactg ccgaagcgcc actcactaaa ctaaccactc catagttta ttaataggat	360
ctggcttgca aacaaaggat gataaacttt gtttatcgct gggagatggg ttggtaacaa	420
aggatgataa actatgttta tcgctggag atgggttaat aacaaaaat gatgtactat	480
gtgccaaact aggacatggc cttgtgttg actcttccaa tgctatcacc atagaaaaca	540
acacccctgtg gacaggcgca aaaccaagcg ccaactgtgt aattaaagag ggagaagatt	600
ccccagactg taagctact ttagttctag tgaagaatgg aggactgata aatggataca	660
taacattaat gggagcctca gaatatacta acacccgtt taaaacaatc aagttacaat	720
cgtgtaaac ctgcatttg ataatactgg ccaaattatt acttacctat catcccttaa	780
aagtaacctg aactttaaag acaaccaaaa catggctact ggaaccataa ccagtgc当地	840
aggcttcatg cccagcacca ccgcctatcc atttataaca tacgccactg agaccctaaa	900
tgaagattac atttatggag agtgttacta caaatctacc aatggaaactc tctttccact	960
aaaagttact gtcacactaa acagacgtat gttagttct ggaatggcct atgctatgat	1020
ttttcatggt ctctaaatgc agaggaagcc ccggaaacta ccgaagtcac tctcattacc	1068
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<210> 33
<211> 1062
<212> DNA
<213> Adenovirus 16

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agcccagatg gagttctaac tcttaatgt gttaatccac tcactaccgc cagcggaccc	240
ctccaactta aagttggaag cagtcttaca gtagatacta tcgatgggtc tttggaggaa	300

tctggcttgc aaacaaagga tgataaactt tgtttatcgc tgggagatgg gttggtaaca 360
 aaggatgata aactatgtt atcgctggga gatgggttaa taacaaaaaa tgatgtacta 420
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 atcgatgtaa acctcgatt tgataatact gccaatttttacttacatccctt atcatccctt 720
 aaaagtaacc tgaactttaa agacaaccaa aacatggcta ctggaaccat aaccagtgcc 780
 aaaggcttca tgcccagcac caccgcctat ccatttataa catacgccac tgagacccta 840
 aatgaagatt acatttatgg agagtgttac tacaaatcta ccaatggaac tctctttcca 900
 ctaaaagtta ctgtcacact aaacagacgt atgttagctt ctggaatggc ctatgctatg 960
 aatttttcat ggtctctaaa tgcagaggaa gccccggaaa ctaccgaagt cactctcatt 1020
 acctccccct tcttttttc ttatatcaga gaagatgact ga 1062

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 <212> PRT
 <213> Artificial sequence

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 Tyr Glu Asp Glu Ser Ser Gln His Pro Phe Ile Asn Pro Gly Phe

20 25 30
 Ile Ser Ser Asn Gly Phe Ala Gln Ser Pro Asp Gly Val Leu Thr Leu
 35 40 45

Lys Cys Val Asn Pro Leu Thr Thr Ala Ser Gly Pro Leu Gln Leu Lys
 50 55 60

Val Gly Ser Ser Leu Thr Val Asp Thr Ile Asp Gly Ser Leu Glu Glu
 65 70 75 80

Asn Ile Thr Ala Ala Pro Leu Thr Lys Thr Asn His Ser Ile Gly

85

90

95

Leu Leu Ile Gly Ser Gly Leu Gln Thr Lys Asp Asp Lys Leu Cys Leu
100 105 110

Ser Leu Glu Asp Gly Leu Val Thr Lys Asp Asp Lys Leu Cys Leu Ser
115 120 125

Leu Gly Asp Gly Leu Ile Thr Lys Asn Asp Val Leu Cys Ala Lys Leu
130 135 140

Gly His Gly Leu Val Phe Asp Ser Ser Asn Ala Ile Thr Ile Glu Asn
145 150 155 160

Asn Thr Leu Trp Thr Gly Ala Lys Pro Ser Ala Asn Cys Val Ile Lys
165 170 175

Glu Gly Glu Asp Ser Pro Asp Cys Lys Leu Thr Leu Val Leu Val Lys
180 185 190

Asn Gly Gly Leu Ile Asn Gly Tyr Ile Thr Leu Met Gly Ala Ser Glu
195 200 205

Tyr Thr Asn Thr Leu Phe Lys Asn Asn Gln Val Thr Ile Asp Val Asn
210 215 220

Leu Ala Phe Asp Asn Thr Gly Gln Ile Ile Thr Tyr Leu Ser Ser Leu
225 230 235 240

Lys Ser Asn Leu Asn Phe Lys Asp Asn Gln Asn Met Ala Thr Gly Thr
245 250 255

Ile Thr Ser Ala Lys Gly Phe Met Pro Ser Thr Thr Ala Tyr Pro Phe
260 265 270

Ile Thr Tyr Ala Thr Glu Thr Leu Asn Glu Asp Tyr Ile Tyr Gly Glu
275 280 285

Cys Tyr Tyr Lys Ser Thr Asn Gly Thr Leu Phe Pro Leu Lys Val Thr
290 295 300

Val Thr Leu Asn Arg Arg Met Leu Ala Ser Gly Met Ala Tyr Ala Met
305 310 315 320

Asn Phe Ser Trp Ser Leu Asn Ala Glu Glu Ala Pro Glu Thr Thr Glu
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Val Thr Leu Ile Thr Ser Pro Phe Phe Ser Tyr Ile Arg Glu Asp
340 345 350

Asp

<210> 35
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<213> Adenovirus Ad16

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Tyr Glu Asp Glu Ser Ser Gln His Pro Phe Ile Asn Pro Gly Phe

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Ile Ser Ser Asn Gly Phe Ala Gln Ser Pro Asp Gly Val Leu Thr Leu
35 40 45

Lys Cys Val Asn Pro Leu Thr Thr Ala Ser Gly Pro Leu Gln Leu Lys
50 55 60

65 70 75 80
Val Gly Ser Ser Leu Thr Val Asp Thr Ile Asp Gly Ser Leu Glu
Lys Cys Val Asn Pro Leu Thr Thr Ala Ser Gly Pro Leu Gln Leu Lys

85 90 95
Asn Ile Thr Ala Ala Ala Pro Leu Thr Lys Thr Asn His Ser Ile Gly

100 105 110
Leu Leu Ile Gly Ser Gly Leu Gln Thr Lys Asp Asp Lys Leu Cys Leu

115 120 125
Ser Leu Gly Asp Gly Leu Val Thr Lys Asp Asp Lys Leu Cys Leu Ser

130 135 140
Leu Gly Asp Gly Leu Ile Thr Lys Asn Asp Val Leu Cys Ala Lys Leu

145 150 155 160
Gly His Gly Leu Val Phe Asp Ser Ser Asn Ala Ile Thr Ile Glu Asn

Asn Thr Leu Trp Thr Gly Ala Lys Pro Ser Ala Asn Cys Val Ile Lys
165 170 175

Glu Gly Glu Asp Ser Pro Asp Cys Lys Leu Thr Leu Val Leu Val Lys
180 185 190

Asn Gly Gly Leu Ile Asn Gly Tyr Ile Thr Leu Met Gly Ala Ser Glu
195 200 205

Tyr Thr Asn Thr Leu Phe Lys Asn Asn Gln Val Thr Ile Asp Val Asn
210 215 220

Leu Ala Phe Asp Asn Thr Gly Gln Ile Ile Thr Tyr Leu Ser Ser Leu
225 230 235 240

Lys Ser Asn Leu Asn Phe Lys Asp Asn Gln Asn Met Ala Thr Gly Thr
245 250 255

Ile Thr Ser Ala Lys Gly Phe Met Pro Ser Thr Thr Ala Tyr Pro Phe
260 265 270

Ile Thr Tyr Ala Thr Glu Thr Leu Asn Glu Asp Tyr Ile Tyr Gly Glu
275 280 285

Cys Tyr Tyr Lys Ser Thr Asn Gly Thr Leu Phe Pro Leu Lys Val Thr
290 295 300

Val Thr Leu Asn Arg Arg Met Leu Ala Ser Gly Met Ala Tyr Ala Met
305 310 315 320

Asn Phe Ser Trp Ser Leu Asn Ala Glu Glu Ala Pro Glu Thr Thr Glu
325 330 335

Val Thr Leu Ile Thr Ser Pro Phe Phe Ser Tyr Ile Arg Glu Asp
340 345 350

Asp

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<223> Chemically synthesized Primer NY-DOWN

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<212> DNA
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180
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240
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300
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